```
# Program: needle
# Rundate: Mon Sep 11 20:02:59 2006
# Align format: srspair
# Report_file: /ebi/extserv/old-work/needle-20060911-20025882318727.output
#
# Aligned sequences: 2
# 1: SEQ 179
# 2: G kaustophilus holB
# Matrix: EBLOSUM62
# Gap penalty: 10.0
# Extend_penalty: 0.5
# Length: 987
           754/987 (76.4%)
# Identity:
           754/987 (76.4%)
# Similarity:
           230/987 (23.3%)
# Gaps:
# Score: 4576.0
#
#-----
             1 atgcgatgggaacagctagcgaaacgccagccggtggtggcgaaaatgct
                                                     50
SEQ_179
               50
G kaustophilu
             1 atqcqatgggaacagctagcgaaacgccagccggtggtggcgaaaatgct
                                                     100
            51 gcaaagcggcttggaaaaagggcggatttctcatgcgtacttgtttgagg
SEQ 179
               51 gcaaagcggcttggaaaaagggcggatttctcatgcgtacttgtttgagg
                                                     1.00
G kaustophilu
            101 qqcaqcqqqqacqggcaaaaaagcggccagtttgttgttggcgaaacgt
                                                    150
SEQ_179
               101 ggcagcggggacgggcaaaaaagcggccagtttgttgttggcgaaacgt
                                                    150
G_kaustophilu
            151 ttgttttgtctgtccccaatcggagtttccccgtgtctagagtgccgcaa
                                                    200
SEQ_179
               200
            151 ttgttttgtctgtccccaatcggagtttccccgtgtctagagtgccgcaa
G kaustophilu
            201 ctgccggcgcatcgactccggcaaccaccctgacgtccgggtgatcggcc
                                                    250
SEQ 179
               201 ctgccggcgcatcgactccggcaaccaccctgacgtccgggtgatcagcc
                                                    250
G kaustophilu
            251 cagatggaggatcaatcaaaaaggaacaaatcgaatggctgcagcaagag
                                                    300
SEQ 179
               251 cagacggaggatcaatcaaaaaggaacaaatcgaatggctgcagcaagag
                                                    300
G_kaustophilu
SEQ_179
            301 ttctcqaaaacagcggtcgagtcggataaaaaaatgtacatcgttgagca
                                                    350
               301 ttctcgaaaacagcggtcgagtcggataaaaaaatgtacatcgttgagca
                                                    350
G kaustophilu
                                                    400
            351 cgccgatcaaatgacgacaagcgctgccaacagccttctgaaatttttgg
SEQ 179
               351 cgccgatcaaatgacgacaagcgctgccaacagccttctgaaatttttgg
                                                    400
G kaustophilu
                                                    450
            SEQ 179
              450
G kaustophilu
           451 cgcctgctagggacgatcgtttcccgctgtcaagtgctttcgttccggcc
                                                    500
SEQ_179
              500
G kaustophilu
           451 egectgetagggaegategttteeegetgteaagtgetttegtteeggee
```

SEQ_179	gttgccgccggcagagctcgcccagggacttgtcgaggagcacgtgccgt		550
G_kaustophilu	501 gttgccgccggcagagctcgctcagggacttgtcgaggag	cacgtgccgt	550
SEQ_179	551 tgccgttggcgctgttggctgcccatttgacaaacagctt	:cgaggaagca	600
G_kaustophilu	551 tgccgttggcgctgttggctgcccatttgacaaacagctt		600
SEQ_179	601 ctggcgcttgccaaagatagttggtttgccgaggcgcgaa		650
G_kaustophilu		ctggcgcttgccaaagatagttggtttgccgaggcgcgaacattagtgct	
SEQ_179	651 acaatggtatgagatgctgggcaagccggagctgcagctt	ttgtttttca	700
G_kaustophilu	651 acaatggtatgagatgctgggcaagccggagctgcagctt		700
SEQ_179	701 tccacgaccgcttgtttccgcattttttggaaagccatca	gettgaeett	750
G_kaustophilu	701 tecaegacegettgttteegeattttttggaaageeatea	tccacgaccgcttgtttccgcattttttggaaagccatcagcttgacctt	
SEQ_179	751 ggacttg		757
G_kaustophilu	751 ggacttgatttgcttttatatttataccgcgatttgttgc	atattcaagc	800
SEQ_179	758		757
G_kaustophilu	801 cgggcagatggacggcgtgttataccgcgatcagctggac	egeetgeaac	850
SEQ_179	758		757
G_kaustophilu	851 gatgggcgcttgcttgcccgcagcggcggattttggctgg	catggaagcg	900
SEQ_179	758		757
G_kaustophilu	901 attttacaagcgaaaacgcgtttaaatacaacgaatatga	gcacggcgtt	950
SEQ_179	758	757	
G_kaustophilu	951 gcttgttgagcagctcgttcttcaattaaagcggtaa	987	

₹.